AUG 1 > 1997

Docket No. 1020

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Patent Application

Group: 1807

A. Ryncarz

Examiner: B. Sisson

Serial Number: 08/791,240

Filed: 01/30/1997

For:

Positive Controls In Polynucleotide Amplification -

COMMUNICATION

August 15, 1997

RECE!

SEP 1 0 1997

Honorable Commissioner of Patents and Trademarks Washington, D.C. 20231

GROUi

Sir:

This is in response to the communication from the Examiner mailed July 29, 1997 in this application.

Enclosed are a replacement diskette and a copy of the Raw Sequence Listing Error Report.

I hereby certify that no data has been changed on this diskette except that the identified errors have been corrected and the serial number has been entered.



Should the Examiner believe that a telephone call to the undersigned would narrow any outstanding issues or favorably advance the prosecution of this application he is respectfully invited to call at the number indicated below.

Respectfully Submitted,

Barbara V. Maurer (Reg. No. 31,278)

Attorney for Applicant(s)
Behring Diagnostics Inc.
3403 Yerba Buena Road
San Jose, CA 95161-9013

(908) 231-4079

CERTIFICATE OF MAILING (37 CFR 1.8a)

I hereby certify that this paper (along with any paper referred to as being attached or enclosed) is being deposited with the United States Postal Service on the date shown below with sufficient postage as first class mail in an envelope addressed to the: the Commissioner of Patents and Trademarks, Washington, D.C. 20231

BARBARA V. MAURER
(Type or print name of person mailing paper)

Date: August 15, 1997

(Signature or person maining paper)

Encl(s) 1 Diskette

Copy of Raw Sequence Listing Error Report

2002

RAW SEQUENCE LISTING PATENT APPLICATION US/08/791,240 DATE: 04/14/97 TIME: 07:46:15

INPUT SET: S16984.raw

This Raw Listing contains the General
Information Section and those Sequences
One Not Comply
Needled Little SEQUENCE LISTING 2 General Information: 3 (i) APPLICANT: Ryncarz, Alexander J. 4 (ii) TITLE OF INVENTION: Positive Controls in Polynucleotide 5 6 Amplification 7 8 (iii) NUMBER OF SEQUENCES: 3 10 11 12 (iv) CORRESPONDENCE ADDRESS: 13 14 Jerome Rosenstock, Esq. 15 ADDRESSEE: (A) 16 Rt. 202-206 North/P.O. Box 2500 17 STREET: (B) 18 Somerville 19 CITY: (C) 20 New Jersey 21 STATE: (D) 23 U.S.A. COUNTRY: (E) 24 08876-1258 25 ZIP: (F) 26 27 COMPUTER READABLE FORM: 28 (V) MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE 29 30 31 DELL 486 32 COMPUTER: (B) 33 WINDOWS 3.1 OPERATING SYSTEM: 34 35 WORDPERFECT 6.0a 36 SOFTWARE: (D) 37 (vi) CURRENT APPLICATION DATA: 38 APPLICATION NUMBER: Not yet Assigned (A) Japuary 30, 1997 FILING DATE: (B) 43 CLASSIFICATION: 44 (C) 45

RAW SEQUENCE LISTING PATENT APPLICATION US/08/791,240

DATE: 04/14/97 TIME: 07:46:18

INPUT SET: S16984.raw

46 47 48

(vii) PRIOR APPLICATION DATA:

49 **5**0

(A) APPLICATION NUMBER:

51 52

(B) FILING DATE:

53 54

(viii) ATTORNEY/AGENT INFORMATION:

55 56 57

(A) NAME:

Jerome Rosenstock, Esq.

58 59

(B) REGISTRATION NUMBER:

25,456

60 61

(C) REFERENCE/DOCKET NUMBER: BDI-1020

62 63

(ix) TELECOMMUNICATION INFORMATION:

64 65

(A) TELEPHONE: (908) 231-2125

66 67

(B) TELEFAX: (908) 231-4919

68 69

ERRORED SEQUENCES FOLLOW:

70 (2) INFORMATION FOR SEQ ID NO:1: 71 72 (i) SEQUENCE CHARACTERISTICS: 73

74 75 76 (A) LENCTH: 25 base pairs

(B) TYPE:

nucleic acid

77 78 79

(C) STRANDEDNESS: single

80

(D) TOPOLOGY: linear

81 82

(ii) MOLECULE TYPE:

83 84

(111) HYPOTHETICAL: NO

85

(iv) ANTI-SENSE:

86 87

(v) FRAGMENT TYPE: C-terminal

88 89 90

(vi) ORIGINAL SOURCE:

91 92

(A) ORGANISM:

see ned poqu

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RAW SEQUENCE LISTING PATENT APPLICATION US/08/791,240

DATE: 04/14/97 TIME: 07:46:21

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INPUT SET: S16984.raw
  93
  94
                  (B) STRAIN:
  95
  96
  97
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
  98
  99
       CGACTCACTA TAGGGCGAAT TGGGC
100
       per 1.822 of Seguese Rules, enumerate bases at the night margin,
per line, giving the cumulative to
101
102
103
       (2) INFORMATION FOR SEQ ID NO:2:
104
105
            (i) SEQUENCE CHARACTERISTICS:
106
107
                 (A) LENGTH:
                                 25 base pairs
108
109
                                nucleic acid
                 (B) TYPE:
110
111
                 (C) STRANDEDNESS: single
112
113
                 (D) TOPOLOGY: linear
114
115
            (ii) MOLECULE TYPE:
116
117
            (iii) HYPOTHETICAL: NO
118
119
            (iv) ANTI-SENSE:
                                 NO
120
121
            (V) FRAGMENT TYPE: C-terminal
122
123
            (vi) ORIGINAL SOURCE:
124
125
                 (A) ORGANISM:
126
127
                 (B) STRAIN:
128
129
           (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: ) They aren't permitted.
130
131
      CGACTCACTA TAGGGCGAAT TGGFGGC
132
133
134
135
136
137
138
139
      (2) INFORMATION FOR SEQ ID NO:3:
140
141
142
           (i) SEQUENCE CHARACTERISTICS:
```





DATE: 04/14/97 TIME: 07:46:25

	1	NPUT SET: S16984.raw
144	(A) LENGTH: 24 base pairs	<u> </u>
145	• • • • • • • •	171
146	(B) TYPE: nucleic acid	CF3
147		له شهر حو
148	(C) STRANDEDNESS: single	
149		
150	(D) TOPOLOGY: linear	
151		
152	(ii) MOLECULE TYPE:	
153		E STATE OF THE STA
154	(111) HYPOTHETICAL: NO	
155	(1x) ANTH-SENSE: NO	CARAMATA CAR
156	(1V) ANTI-SENSE: NO	· -
157		
158	(V) FRAGMENT TYPE: C-terminal	
159	A ' ANTERWAL COMMON.	
160	(Vi) ORIGINAL SOURCE:	• • • • • • • • • • • • • • • • • • •
161	AN ORGANISM.	
162	(A) ORGANISM:	•
163	(B) STRAIN:	
164	(B) SIRAIN:	•
165		, sest
166	(Ri) SEQUENCE DESCRIPTION: SEQ ID NO: 3:	Visut 24
167	(XT) PETABLED PROGNET PROMISE TO THE	24
168	CATTAGGCAC CCCAGGCTTT ACAC	24
169	CATTAGGCAC CCCAGGOIII	

RAW SEQUENCE LISTING PATENT APPLICATION US/08/791,240





SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/08/791,240

DATE: 04/14/97 TIME: 07:46:28

INPUT SET: S16984.raw

Line	Error	Original Text
41 74 107 144	Wrong application Scrial Number Entered (25) and Calc. Seq. Length (0) differ Entered (25) and Calc. Seq. Length (0) differ Entered (24) and Calc. Seq. Length (0) differ	 (A) APPLICATION NUMBER: Not yet Assigned (A) LENGTH: 25 base pairs (A) LENGTH: 25 base pairs (A) LENGTH: 24 base pairs

DEG ASSESSED COPY